HS prok arch filtered Differential expression AvH

All ✓ records per page		Search all columns:			
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ID	Image		logFC		Adjus p-Val⊧
k141_181201_25_transcriptional_regulators	ф ф	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181201_25_transcripti			3.36e
k141_181201_19_Domain_of_unknown_function_DUF302	• • • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181201_19_Domain_o	-7.80	4.95e-09	7.80e
k141_372642_30	• • • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_372642_30pdf)	-6.74	4.98e-08	5.40e
k141_181201_18	ф Ф	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181201_18pdf)	-6.67	2.22e-11	5.24e
k141_181201_23	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181201_23pdf)	-5.71	1.51e-03	3.40e
k141_92599_11_COG1522_Transcriptional_regulators	• ••	$(figures HS_prok_arch_RNAs eq_analysis_with_DES eq2_longer_names_AvH/boxplot.k141_92599_11_COG1522\$	-5.71	3.42e-04	8.50e
k141_92599_5_helix_turn_helix_ASNC_type	ф ш о <u>«тө</u> »	$(figures HS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_92599_5_helix_turn_h$	-5.60	4.10e-05	1.62e
k141_181201_20_transcriptional_regulators		(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181201_20_transcripti	-5.28	5.63e-03	9.73e
k141_92599_3_Lrp_AsnC_ligand_binding_domain		(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_92599_3_trp_AsnC_li	-5.27	5.71e-08	5.40e
k141_92599_12_COG1522_Transcriptional_regulators		(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_92599_12_COG1522	-5.15	1.65e-03	3.55e
k141_96470_27_CBS_domain		(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_96470_27_CBS_doma	-4.85	6.94e-06	3.30e
k141_447824_6	ф 613	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_447824_6pdf)	-4.61	8.16e-12	3.86e
k141_98648_15_PFAM_bluetype_1copper_domain_pro	•	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_98648_15_PFAM_blue	-4.40	1.16e-06	9.16e
k141_392539_6_COG0589_Universal_stress_protein_Usp	• • • • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_392539_6_COG0589	-4.37	2.38e-06	1.60e
k141_326599_17_Universal_stress_protein	• • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_326599_17_Universal	-4.34	6.97e-06	3.30e
k141_92599_10_COG1522_Transcriptional_regulators	•	$(figures HS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_92599_10_COG1522\$	-3.93	1.23e-04	3.88e
k141_326599_21_CBS_domain	•	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_326599_21_CBS_dom	-3.84	9.96e-05	3.36e
k141_331008_5_Nitroreductase_436308_Nmar_0360	ф ф-ф	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_331008_5_Nitroreduct	-3.71	2.22e-03	4.56e
k141_440873_29_Belongs_to_the_small_heat_shock_pro	0 0 0 0 0 0	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_440873_29_Belongs_t	-3.69	4.42e-04	1.05e
k141_440873_33_Stabilizes_TBP_binding_to_an_archae		(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_440873_33_Stabilizes	-3.65	3.93e-06	2.32e
k141_15183_34	•	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_15183_34pdf)	-2.90	3.62e-05	1.56e
k141_181885_20_COG1522_Transcriptional_regulators	6 10	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_181885_20_COG1522	-2.70	2.29e-04	6.73e
k141_381857_3_PFAM_Integrase_core_domain	• • • • • • • • • • • • • • • • • • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_381857_3_PFAM_Inte	-2.64	2.86e-04	7.52e
k141_76157_13_Belongs_to_the_P_IIprotein_family	• • • •	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_76157_13_Belongs_to	-2.63	4.44e-03	8.07e
k141_93077_13	ф ф «о зы	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_93077_13pdf)	-2.44	2.42e-04	6.73e

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ID	Image	logFC		Adjus p-Val
k141_302809_9_Short-chain_dehydrogenase_reductase_	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_302809_9_Short-chain	-2.09	2.37e-03	4.68e
k141_198002_28	* (figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_198002_28pdf)	-1.63	3.62e-03	6.85e
k141_339108_19	(figuresHS_prok_arch_RNAseq_analysis_with_DESeq2_longer_names_AvH/boxplot.k141_339108_19pdf)	1.88	5.76e-03	9.73e
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